### **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/5/6,5/7Source: 10/5/6,5/7Date Processed by STIC: 12-10-0

# ENTERED



PCT

#### RAW SEQUENCE LISTING

DATE: 12/10/2004 TIME: 14:26:34

PATENT APPLICATION: US/10/516,517

Input Set : A:\Q85059.txt

Output Set: N:\CRF4\12102004\J516517.raw

```
3 <110> APPLICANT: Yamanouchi Pharmaceutical Co., Ltd.
      4
              Kentaro ENJO
      5
              Sadao KUROMITSU
      7 <120> TITLE OF INVENTION: Method for screening an agent for treating renal failure
      9 <130> FILE REFERENCE: Q85059
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/516,517
C--> 11 <141> CURRENT FILING DATE: 2004-12-02
     11 <150> PRIOR APPLICATION NUMBER: PCT/JP03/012967
     12 <151> PRIOR FILING DATE: 2003-10-09
     14 <150> PRIOR APPLICATION NUMBER: JP2002-298958
     15 <151> PRIOR FILING DATE: 2002-10-11
    17 <160> NUMBER OF SEQ ID NOS: 20
     19 <170> SOFTWARE: PatentIn version 3.1
     21 <210> SEQ ID NO: 1
     22 <211> LENGTH: 993
     23 <212> TYPE: DNA
     24 <213> ORGANISM: Homo sapiens
     26 <220> FEATURE:
     27 <221> NAME/KEY: CDS
     28 <222> LOCATION: (1)..(990)
     29 <223> OTHER INFORMATION: inventor: Enjo, Kentaro; Kuromitsu, Sadao
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                                                                               48
     34 Met Ala Trp Asn Ala Thr Cys Lys Asn Trp Leu Ala Ala Glu Ala Ala
                                            10
     37 ctg gaa aag tac tac ctt tcc att ttt tat ggg att gag ttc gtt gtg
                                                                               96
    38 Leu Glu Lys Tyr Tyr Leu Ser Ile Phe Tyr Gly Ile Glu Phe Val Val
                    20
    41 gga gtc ctt gga aat acc att gtt gtt tac ggc tac atc ttc tct ctq
    42 Gly Val Leu Gly Asn Thr Ile Val Val Tyr Gly Tyr Ile Phe Ser Leu
    45 aag aac tgg aac agc agt aat att tat ctc ttt aac ctc tct gtc tct
                                                                              192
     46 Lys Asn Trp Asn Ser Ser Asn Ile Tyr Leu Phe Asn Leu Ser Val Ser
    47
            50
    49 gac tta gct ttt ctg tgc acc ctc ccc atg ctg ata agg agt tat gcc
                                                                              240
    50 Asp Leu Ala Phe Leu Cys Thr Leu Pro Met Leu Ile Arg Ser Tyr Ala
    53 aat gga aac tgg ata tat gga gac gtg ctc tgc ata agc aac cga tat
                                                                              288
    54 Asn Gly Asn Trp Ile Tyr Gly Asp Val Leu Cys Ile Ser Asn Arg Tyr
                        85
                                            90
    57 gtg ctt cat gcc aac ctc tat acc agc att ctc ttt ctc act ttt atc
                                                                              336
    58 Val Leu His Ala Asn Leu Tyr Thr Ser Ile Leu Phe Leu Thr Phe Ile
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100

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## RAW SEQUENCE LISTING DATE: 12/10/2004 PATENT APPLICATION: US/10/516,517 TIME: 14:26:34

Input Set : A:\Q85059.txt

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61 agc ata gat co				
62 Ser Ile Asp Ar	g Tyr Leu Ile		_	lu His Leu
63 115	•	120	125	
65 ctg caa aag aa				
66 Leu Gln Lys Ly		Ile Leu Ile		le Trp Val
67 130	135		140	
69 tta gta acc tt				
70 Leu Val Thr Le		Pro Ile Leu	Pro Leu Ile As	sn Pro Val
71 145	150		155	160
73 ata act gac aa				
74 Ile Thr Asp As			Phe Ala Ser Se	
75	165	170		175
77 ccc aac tac aa				
78 Pro Asn Tyr As		_		<del>-</del>
79 18		185	19	
81 ctt att cct ct				
82 Leu Ile Pro Le	u Phe Val Met			le Ala Leu
83 195		200	205	
85 ttc cta aag ca				
86 Phe Leu Lys Gl		Gln Val Ala		co Leu Glu
87 210	215		220	•
89 aag cct ctc aa				
90 Lys Pro Leu As		Met Ala Val		
91 225	230		235	240
93 ttt aca ccc ta				
94 Phe Thr Pro Ty			Arg Ile Ala Se	
95	245	250		255
97 ggg agt tgg aa				
98 Gly Ser Trp Ly 99 26				
		265	2'	
101 tac att gtg a				
102 Tyr Ile Val T 103 275	iii Aig Aia Le	280	a Ash Ser var . 285	ile Ash Pro
105 gtc ttc tat t	tt att tta aa			etg atg aat 912
106 Val Phe Tyr F	he Leu Leu Gl	u Aen Wie Ph	ayy yac acy (	ley Mot Agn
107 290	29		300	led Met Asii
109 caa ctg aga c				aga tgg gct 960
110 Gln Leu Arg H				
111 305	310	b ber bed in	315	320
113 cat gaa ctc c		r ana naa aa		993
114 His Glu Leu I				993
115	325	33		
118 <210> SEQ ID NO: 2				
119 <211> LENGTH: 330				
120 <212> TYPE: F				
121 <213> ORGANISM: Homo sapiens				
123 <400> SEQUENC				
125 Met Ala Trp A		s Lys Asn Tr	Leu Ala Ala (	Slu Ala Ala
126 1	5	10		15

**RAW SEQUENCE LISTING**PATENT APPLICATION: **US/10/516,517**DATE: 12/10/2004

TIME: 14:26:34

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129 Leu Glu Lys Tyr Tyr Leu Ser Ile Phe Tyr Gly Ile Glu Phe Val Val
     133 Gly Val Leu Gly Asn Thr Ile Val Val Tyr Gly Tyr Ile Phe Ser Leu
     137 Lys Asn Trp Asn Ser Ser Asn Ile Tyr Leu Phe Asn Leu Ser Val Ser
                                55
     141 Asp Leu Ala Phe Leu Cys Thr Leu Pro Met Leu Ile Arg Ser Tyr Ala
     145 Asn Gly Asn Trp Ile Tyr Gly Asp Val Leu Cys Ile Ser Asn Arg Tyr
                        85
                                            90
     149 Val Leu His Ala Asn Leu Tyr Thr Ser Ile Leu Phe Leu Thr Phe Ile
                                        105
     153 Ser Ile Asp Arg Tyr Leu Ile Ile Lys Tyr Pro Phe Arg Glu His Leu
                                    120
     157 Leu Gln Lys Lys Glu Phe Ala Ile Leu Ile Ser Leu Ala Ile Trp Val
                                135
                                                    140
     161 Leu Val Thr Leu Glu Leu Leu Pro Ile Leu Pro Leu Ile Asn Pro Val
                            150
                                                155
     165 Ile Thr Asp Asn Gly Thr Thr Cys Asn Asp Phe Ala Ser Ser Gly Asp
                        165
                                            170
     169 Pro Asn Tyr Asn Leu Ile Tyr Ser Met Cys Leu Thr Leu Leu Gly Phe
     170
                                        185
    173 Leu Ile Pro Leu Phe Val Met Cys Phe Phe Tyr Tyr Lys Ile Ala Leu
               195
                                    200
                                                        205
    177 Phe Leu Lys Gln Arg Asn Arg Gln Val Ala Thr Ala Leu Pro Leu Glu
                                215
     181 Lys Pro Leu Asn Leu Val Ile Met Ala Val Val Ile Phe Ser Val Leu
                            230
                                                235
    185 Phe Thr Pro Tyr His Val Met Arg Asn Val Arg Ile Ala Ser Arg Leu
    189 Gly Ser Trp Lys Gln Tyr Gln Cys Thr Gln Val Val Ile Asn Ser Phe
                    260
                                        265
    193 Tyr Ile Val Thr Arg Ala Leu Gly Phe Leu Asn Ser Val Ile Asn Pro
    194 275
                                    280
    197 Val Phe Tyr Phe Leu Leu Gly Asp His Phe Arg Asp Met Leu Met Asn
                                295
                                                    300
    201 Gln Leu Arg His Asn Phe Lys Ser Leu Thr Ser Phe Ser Arg Trp Ala
    202 305
                         310
                                                315
    205 His Glu Leu Leu Ser Phe Arg Glu Lys
                        325
    209 <210> SEQ ID NO: 3
    210 <211> LENGTH: 29
    211 <212> TYPE: DNA
    212 <213> ORGANISM: Artificial Sequence
    214 <220> FEATURE:
    215 <223> OTHER INFORMATION: Description of Artificial Sequence:an artificially
systhesized
    216
            primer sequence
    218 <400> SEQUENCE: 3
    219 ggtctagaat ggcatggaat gcaacttgc
                                                                              29
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### RAW SEQUENCE LISTING

DATE: 12/10/2004 TIME: 14:26:34

Input Set : A:\Q85059.txt

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PATENT APPLICATION: US/10/516,517

222 <210> SEQ ID NO: 4 223 <211> LENGTH: 32 224 <212> TYPE: DNA 225 <213> ORGANISM: Artificial Sequence 227 <220> FEATURE: 228 <223> OTHER INFORMATION: Description of Artificial Sequence: an artificially systhesized 229 primer sequence 231 <400> SEQUENCE: 4 232 ggtctagatt atcacttttc tctqaatqaa aq 32 235 <210> SEQ ID NO: 5 236 <211> LENGTH: 19 237 <212> TYPE: DNA 238 <213> ORGANISM: Homo sapiens 240 <400> SEQUENCE: 5 241 tcaggctgca tgttccttg 19 244 <210> SEQ ID NO: 6 245 <211> LENGTH: 19 246 <212> TYPE: DNA 247 <213> ORGANISM: Homo sapiens 249 <400> SEQUENCE: 6 250 tecteteage ggggaagag 19 253 <210> SEQ ID NO: 7 . 254 <211> LENGTH: 21 255 <212> TYPE: DNA 256 <213> ORGANISM: Homo sapiens 258 <400> SEQUENCE: 7 259 gacgtgtccc atagtgtttc c 21 262 <210> SEQ ID NO: 8 263 <211> LENGTH: 18 264 <212> TYPE: DNA 265 <213> ORGANISM: Homo sapiens 267 <400> SEQUENCE: 8 268 teetgetgee gtggtget 18 271 <210> SEO ID NO: 9 272 <211> LENGTH: 27 273 <212> TYPE: DNA 274 <213> ORGANISM: Artificial Sequence 276 <220> FEATURE: 277 <223> OTHER INFORMATION: Description of Artificial Sequence:an artificially systhesized 278 primer sequence 280 <400> SEQUENCE: 9 281 ggggtacctc aggctgcatg ttccttg 27 284 <210> SEQ ID NO: 10 285 <211> LENGTH: 28 286 <212> TYPE: DNA 287 <213> ORGANISM: Artificial Sequence 289 <220> FEATURE: 290 <223> OTHER INFORMATION: Description of Artificial Sequence: an artificially systhesized

primer sequence

291

**RAW SEQUENCE LISTING**PATENT APPLICATION: **US/10/516,517**DATE: 12/10/2004

TIME: 14:26:34

Input Set : A:\Q85059.txt

Output Set: N:\CRF4\12102004\J516517.raw

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    294 ggaagatett eeteteageg gggaagag
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     298 <211> LENGTH: 29
    299 <212> TYPE: DNA
     300 <213> ORGANISM: Artificial Sequence
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     303 <223> OTHER INFORMATION: Description of Artificial Sequence: an artificially
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     304
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     306 <400> SEQUENCE: 11
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     311 <211> LENGTH: 27
     312 <212> TYPE: DNA
     313 <213> ORGANISM: Artificial Sequence
     315 <220> FEATURE:
     316 <223> OTHER INFORMATION: Desctiption of Artificial Sequence: an artificially
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     317
              primer sequence
     319 <400> SEQUENCE: 12
     320 ggaagatett cetgetgeeg tggtget
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     324 <211> LENGTH: 1159
     325 <212> TYPE: DNA
     326 <213> ORGANISM: Homo sapiens
     328 <400> SEQUENCE: 13
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     331 agtcactaaa aatatattat tcactgtcaa tcttagttta tatccagata caacagggta
                                                                               120
     333 cactgetett gtaatggaat cagacttett attttaacaa gacaaaccaa atecaateca
                                                                               180
     335 catttgaaga ttataggttt taatataaga aaatgcactc atttctcaaa gaccctagtg
                                                                               240
     337 aagetgtgtt taaatgetee taggtgaace ceetttgeat ceeagtgtte ceaecetgae
     339 acccagagee cetacetace caacacagaa teatttgete tgatagaaca atggateeet
                                                                               360
     341 ttttctggaa acattgatgg ccactcctcc cttgtccttg cctatataaa actcctacat
                                                                               420
     343 atattaagag aaaactaagc aagagttttg gaaatctgcc ccaggagact gcatcctgag
                                                                               480
     345 tcacacgegt ctttgttctc tttcttgtcc caaaaccgtt acctcaagtg acaaatgatc
     347 aaateteaaa tatagaatte agggttttae aggtaggeat ettgaggatt teaaatggtt
                                                                               600
     349 aaaagcaact cactcetttt ctactetttg gagagtttca agageetata geetetaaaa
                                                                               660
     351 cgcaaatcat tgctaagggt tgggggggag aaaccttttc gaatttttta ggaattcctg
                                                                               720
     353 ctgtttgcct cttcagctac ctacttccta aaaaggatgt atgtcagtgg acagaacagg
     355 gcaaacttat tcgaaaaaga aataagaaat aattgccagt gtgtttataa atgatatgaa
                                                                               840
     357 tcaggagtgg tgcgaagagg atagggaaaa aaaaattcta tttggtgctg gaaatactgc
                                                                               900
     359 getttttttt tteetttttt tttttttetg tgagetggag tgtgecaget tttteagaeg
                                                                               960
     361 gaggaatget gagtgteaag gggteaggat caateeggtg tgagttgatg aggeaggaag
                                                                              1020
     363 gtggggagga atgcgaggaa tgtccctgtt tgtgtaggac tccattcagc tcattggcga
                                                                              1080
     365 geogeggeeg eeeggagegt ataaaageet egggeegeee geeecaaaet cacacaacaa
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     367 ctcttccccq ctqaqaqqa
                                                                              1159
     370 <210> SEQ ID NO: 14
     371 <211> LENGTH: 355
     372 <212> TYPE: DNA
     373 <213> ORGANISM: Homo sapiens
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### VERIFICATION SUMMARY

PATENT APPLICATION: US/10/516,517

DATE: 12/10/2004 TIME: 14:26:35

Input Set : A:\Q85059.txt

Output Set: N:\CRF4\12102004\J516517.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date